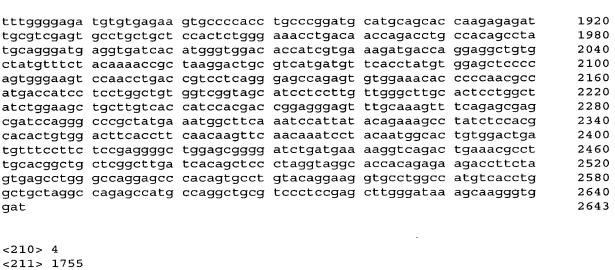
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Pro Tyr Gly Gln Glu Gly Leu Lys Ser Lys Ser Val Gln Leu Asn Gly 425

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Cys Ala Trp Cys Ser Asp Glu Ala Leu Pro Leu Gly Ser Pro Arg Cys

Asp Leu Lys Glu Asn Leu Leu Lys Asp Asn Cys Ala Pro Glu Ser Ile

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Asp Lys Gly Ser Gly Asp Ser Ser Gln Val Thr Gln Val Ser Pro Gln
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Leu Asn Asn Glu Val Ile Pro Gly Leu Lys Ser Cys Met Gly Leu Lys 405 410 415

Asp Ser Leu Ile Val Gln Val Thr Phe Asp Cys Asp Cys Ala Cys Gln 450 455 460

Ala Gln Ala Glu Pro Asn Ser His Arg Cys Asn Asn Gly Asn Gly Thr 465 470 475 480

Tyr Val Cys Gly Leu Cys Glu Cys Ser Pro Gly Tyr Leu Gly Thr Arg 485 490 495

Cys Glu Cys Gln Asp Gly Glu Asn Gln Ser Val Tyr Gln Asn Leu Cys 500 505 510

Arg Glu Ala Glu Gly Lys Pro Leu Cys Ser Gly Arg Gly Asp Cys Ser 515 520 525

Cys Asn Gln Cys Ser Cys Phe Glu Ser Glu Phe Gly Lys Ile Tyr Gly 530 540

Pro Phe Cys Glu Cys Asp Asn Phe Ser Cys Ala Arg Asn Lys Gly Val 545 550 555 560

Leu Cys Ser Gly His Gly Glu Cys His Cys Gly Glu Cys Lys Cys His 565 570 575

Ala Gly Tyr Ile Gly Asp Asn Cys Asn Cys Ser Thr Asp Ile Ser Thr 580 585 590

Cys Arg Gly Arg Asp Gly Gln Ile Cys Ser Glu Arg Gly His Cys Leu 595 600 605

Cys Gly Gln Cys Gln Cys Thr Glu Pro Gly Ala Phe Gly Glu Met Cys 610 620

Glu Lys Cys Pro Thr Cys Pro Asp Ala Cys Ser Thr Lys Arg Asp Cys 625 630 635 640

Val Glu Cys Leu Leu His Ser Gly Lys Pro Asp Asn Gln Thr Cys 645 650 655

His Ser Leu Cys Arg Asp Glu Val Ile Thr Trp Val Asp Thr Ile Val 660 665 670

Lys Asp Asp Gln Glu Ala Val Leu Cys Phe Tyr Lys Thr Ala Lys Asp 675 680 685

Cys Val Met Met Phe Thr Tyr Val Glu Leu Pro Ser Gly Lys Ser Asn 690 695 700

Leu Thr Val Leu Arg Glu Pro Glu Cys Gly Asn Thr Pro Asn Ala Met 705 710 715 720

Thr Ile Leu Leu Ala Val Val Gly Ser Ile Leu Leu Val Gly Leu Ala 725 730 735

Leu Leu Ala Ile Trp Lys Leu Leu Val Thr Ile His Asp Arg Arg Glu 745

Phe Ala Lys Phe Gln Ser Glu Arg Ser Arg Ala Arg Tyr Glu Met Ala 755

Ser Asn Pro Leu Tyr Arg Lys Pro Ile Ser Thr His Thr Val Asp Phe 775

Thr Phe Asn Lys Phe Asn Lys Ser Tyr Asn Gly Thr Val Asp

<210> 9

<211> 315

<212> PRT

<213> Homo sapiens

Met Ala Asn Cys Ser Leu Tyr Arg Ser Cys Gly Asp Cys Leu Leu Ala

Arg Asp Pro Tyr Cys Ala Trp Ser Gly Ser Ser Cys Lys His Val Ser 25

Leu Tyr Gln Pro Gln Leu Ala Thr Arg Pro Trp Ile Gln Asp Ile Glu 35

Gly Ala Ser Ala Lys Asp Leu Cys Ser Ala Ser Ser Val Val Ser Pro

Ser Phe Val Pro Thr Gly Glu Lys Pro Cys Glu Gln Val Gln Phe Gln

Pro Asn Thr Val Asn Thr Leu Ala Cys Pro Leu Leu Ser Asn Leu Ala

Thr Arg Leu Trp Leu Arg Asn Gly Ala Pro Val Asn Ala Ser Ala Ser

Cys His Val Leu Pro Thr Gly Asp Leu Leu Val Gly Thr Gln Gln 120

Leu Gly Glu Phe Gln Cys Trp Ser Leu Glu Glu Gly Phe Gln Gln Leu 130

Val Ala Ser Tyr Cys Pro Glu Val Val Glu Asp Gly Val Ala Asp Gln 155 150

Thr Asp Glu Gly Gly Ser Val Pro Val Ile Ile Ser Thr Ser Arg Val 170 165

Ser Ala Pro Ala Gly Gly Lys Ala Ser Trp Gly Ala Asp Arg Ser Tyr

Trp Lys Glu Phe Leu Val Met Cys Thr Leu Phe Val Leu Ala Val Leu 195

Leu Pro Val Leu Phe Leu Leu Tyr Arg His Arg Asn Ser Met Lys Val

210 215 220

Phe Leu Lys Gln Gly Glu Cys Ala Ser Val His Pro Lys Thr Cys Pro 225 230 235 240

Val Val Leu Pro Pro Glu Thr Arg Pro Leu Asn Gly Leu Gly Pro Pro 245 250 255

Ser Thr Pro Leu Asp His Arg Gly Tyr Gln Ser Leu Ser Asp Ser Pro

Pro Gly Ser Arg Val Phe Thr Glu Ser Glu Lys Arg Pro Leu Ser Ile 275 280 285

Gln Asp Ser Phe Val Glu Val Ser Pro Val Cys Pro Arg Pro Arg Val 290 295 300

Arg Leu Gly Ser Glu Ile Arg Asp Ser Val Val 305 310 315

<210> 10

<211> 375

<212> PRT

<213> Homo sapiens

<400> 10

Met Glu Phe Glu Ile Thr Phe Arg Pro Asp Ser Gly Asp Gly Val Leu
1 5 10 15

Leu Tyr Ser Tyr Asp Thr Gly Ser Lys Asp Phe Leu Ser Ile Asn Leu 20 25 30

Ala Gly Gly His Val Glu Phe Arg Phe Asp Cys Gly Ser Gly Thr Gly 35 40 45

Val Leu Arg Ser Glu Asp Pro Leu Thr Leu Gly Asn Trp His Glu Leu 50 55 60

Arg Val Ser Arg Thr Ala Lys Asn Gly Ile Leu Gln Val Asp Lys Gln 65 70 75 80

Lys Ile Val Glu Gly Met Ala Glu Gly Gly Phe Thr Gln Ile Lys Cys 85 90 95

Asn Thr Asp Ile Phe Ile Gly Gly Val Pro Asn Tyr Asp Asp Val Lys $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110 \hspace{1.5cm}$

Lys Asn Ser Gly Val Leu Lys Pro Phe Ser Gly Ser Ile Gln Lys Ile 115 120 125

Ile Leu Asn Asp Arg Thr Ile His Val Lys His Asp Phe Thr Ser Gly
130 135 140

Val Asn Val Glu Asn Ala Ala His Pro Cys Val Arg Ala Pro Cys Ala 145 150 155 160

His Gly Gly Ser Cys Arg Pro Arg Lys Glu Gly Tyr Asp Cys Asp Cys 165 170 175 Pro Leu Gly Phe Glu Gly Leu His Cys Gln Lys Ala Ile Ile Glu Ala 180 185 190

Ile Glu Ile Pro Gln Phe Ile Gly Arg Ser Tyr Leu Thr Tyr Asp Asn 195 200 205

Pro Asp Ile Leu Lys Arg Val Ser Gly Ser Arg Ser Asn Val Phe Met 210 215 220

Arg Phe Lys Thr Thr Ala Lys Asp Gly Leu Leu Leu Trp Arg Gly Asp 225 230 235 240

Ser Pro Met Arg Pro Asn Ser Asp Phe Ile Ser Leu Gly Leu Arg Asp
245
250
255

Gly Ala Leu Val Phe Ser Tyr Asn Leu Gly Ser Gly Val Ala Ser Ile 260 265 270

Met Val Asn Gly Ser Phe Asn Asp Gly Arg Trp His Arg Val Lys Ala 275 280 285

Val Arg Asp Gly Gln Ser Gly Lys Ile Thr Val Asp Asp Tyr Gly Ala 290 295 300

Arg Thr Gly Lys Ser Pro Gly Met Met Arg Gln Leu Asn Ile Asn Gly 305 310 315 320

Ala Leu Tyr Val Gly Gly Met Lys Glu Ile Ala Leu His Thr Asn Arg 325 330 . 335

Gln Tyr Met Arg Gly Leu Val Gly Cys Ile Ser His Phe Thr Leu Ser 340 345 350

Thr Asp Tyr His Ile Ser Leu Val Glu Asp Ala Val Asp Gly Lys Asn 355 360 365

Ile Asn Thr Cys Gly Ala Lys 370 375

<210> 11

<211> 211

<212> PRT

<213> Homo sapiens

<400> 11

Gln Ile Ser Ala Ala Asp Leu Asp Ser Pro Ala Ser Pro Ile Arg Tyr 1 5 10 15

Ser Ile Leu Pro His Ser Asp Pro Glu Arg Cys Phe Ser Ile Gln Pro 20 25 30

Glu Glu Gly Thr Ile His Thr Ala Ala Pro Leu Asp Arg Glu Ala Arg 35 40 45

Ala Trp His Asn Leu Thr Val Leu Ala Thr Glu Leu Asp Ser Ser Ala
50 55 60

Gln Ala Ser Arg Val Gln Val Ala Ile Gln Thr Leu Asp Lys Asn Asp 65 70 75 80

Asn Ala Pro Gln Leu Ala Glu Pro Tyr Asp Thr Phe Val Cys Asp Ser

Ala Ala Pro Gly Gln Leu Ile Gln Val Ile Arg Ala Leu Asp Arg Asp

Glu Val Gly Asn Ser Ser His Val Ser Phe Gln Gly Pro Leu Gly Pro 120

Asp Ala Asn Phe Thr Val Gln Asp Asn Arg Asp Gly Ser Ala Ser Leu 135

Leu Leu Pro Ser Arg Pro Ala Pro Pro Arg His Ala Pro Tyr Leu Val 155

Pro Ile Glu Leu Trp Asp Trp Gly Gln Pro Ala Leu Ser Ser Thr Ala

Thr Val Thr Val Ser Val Cys Arg Cys Gln Pro Asp Gly Ser Val Ala

Ser Cys Leu Pro Trp Trp Cys Ser Ser Trp Pro Cys Gly Gly Arg Ser

Lys Lys His 210

<210> 12

<211> 439

<212> PRT

<213> Homo sapiens

Gly Asp Arg Arg Pro Leu Pro Val Asp Arg Ala Ala Gly Leu Lys Glu

Lys Thr Leu Ile Leu Leu Asp Val Ser Thr Lys Asn Pro Val Arg Thr

Val Asn Glu Asn Phe Leu Ser Leu Gln Leu Asp Pro Ser Ile Ile His

Asp Gly Trp Leu Asp Phe Leu Ser Ser Lys Arg Leu Val Thr Leu Ala

Arg Gly Leu Ser Pro Ala Phe Leu Arg Phe Gly Gly Lys Arg Thr Asp

Phe Leu Gln Phe Gln Asn Leu Arg Asn Pro Ala Lys Ser Arg Gly Gly . 90

Pro Gly Pro Asp Tyr Tyr Leu Lys Asn Tyr Glu Asp Glu Pro Asn Asn 105

Tyr Arg Thr Met His Gly Arg Ala Val Asn Gly Ser Gln Leu Gly Lys 120

Asp Tyr Ile Gln Leu Lys Ser Leu Leu Gln Pro Ile Arg Ile Tyr Ser

	130						135						1	40					
Arg 145	Ala	Ser	Leu	Τì	/r G	31y :	Pro	Asn	I	le	Gly	Arg 155	J F	ro.	Arg	Lys	Ası	n V	/al 160
Ile	Ala	Leu	Leu	1 As	sp (Sly	Pḥe	Met	L	ys	Val 170	Ala	a C	Bly	Ser	Thr	Va 17	1 <i>1</i> 5	4sp
Ala	Va1	Thr	Trp		ln H	His	Cys	Tyr	1 1	le .85	Asp	Gly	y P	Arg	Val	Val 190	Ly	s \	Val
Met	Asp	Phe		ı L	ys '	Thr	Arg	Leu 200	ı I	eu	Asp	Th	r 1	Leu	Ser 205	Asp	Gl	n	Ile
Arg	Lys 210	Ile	Gli	n L	ys '	Val	Val 215	Asn	ı 1	Chr	Tyr	Th	r i	Pro 220	Gly	Lys	Ьу	ទ	Ile
Trp 225	Leu	Glu	Gl;	y V	al	Val 230	Thr	Thi	<i>:</i>	Ser	Ala	Gl 23	У ⁽ 5	Gly	Thr	Asn	. As	sn	Leu 240
Ser	Asp	Ser	ту	r A	la 245	Ala	Gly	Phe	e 1	Leu	Trp 250	Le	u	Asn	Thr	Leu	G] 25	ly 55	Met
Leu	Ala	Ası	n Gl 26		ly	Ile	Asp	Va:	1 :	Val 265	Ιlε	e Ar	g	His	Ser	Phe 270	e P]	ne	Asp
His	Gly	7 Ty:		n F	His	Leu	Va]	As ₁	р 0	Gln	Asr	n Ph	ıe	Asn	Pro 285	Leu ;	1 P	ro	Asp
Tyr	Trp 290		u S∈	er l	Leu	Leu	Ту: 29!	c Ly	s	Arg	Lev	ı I	le	Gly 300	Pro	b Lys	s V	al	Leu
Ala 305		l Hi	s Va	al A	Ala	Gly 310	Le	u Gl	n	Arg	Ly	3 P	ro 15	Arg	Pro	o Gly	y A	rg	Val 320
Ile	e Ar	g As	p Ly	/s	Leu 325	Arg	11	е Ту	r	Ala	Hi 33	s C 0	ys	Thr	Ası	n Hi	s H 3	is 35	Asn
Hi	s As	n Ty		al 40	Arg	Gly	, Se	r Il	Le	Th:	c Le	u P	he	116	e Il	e As 35	n I O	eu	His
Ar	g Se	r Ar 35		ys	Lys	Ile	e Ly	s Le	eu 50	Ala	a Gl	уТ	hr	Lei	ı Ar 36	g As 5	рΙ	ъys	Leu
Va	1 Hi 37		ln T	yr	Leu	Le	u Gl 37	.n Pi	ro	Ту	r Gl	y G	ln	Gli 38	u Gl O	y Le	eu I	ГÀа	Ser
L у 38		er Va	al G	ln	Leu	1 As: 39	n G] 0	y G	ln	Pr	o Le	u V	7al 895	Me	t Va	ıl As	sp i	Asp	Gly 400
Th	ır Le	eu P	ro G	lu	Leu 405	ı Ly	s Pi	co A	rg	Pr	o Le	eu <i>I</i> LO	٩r٥	g Al	a G]	y Aı	cg '	Th:	Leu 5
Va	al I	le P		Pro 120	Va:	l Th	r Me	et G	ly	Ph 42	e Pl	ne V	√a]	l Va	.1 Ly	/s As	sn 30	Va:	l Asr

Ala Leu Ala Cys Arg Tyr Arg



<212> PRT

<213> Homo sapiens

<400> 13

Met Arg Val Leu Cys Ala Phe Pro Glu Ala Met Pro Ser Ser Asn Ser 1 10 15

Arg Pro Pro Ala Cys Leu Ala Pro Gly Ala Leu Tyr Leu Ala Leu Leu 20 25 30

Leu His Leu Ser Leu Ser Ser Gln Ala Gly Asp Arg Arg Pro Leu Pro
35 40 45

Val Asp Arg Ala Ala Gly Leu Lys Glu Lys Thr Leu Ile Leu Leu Asp 50 55 60

Val Ser Thr Lys Asn Pro Val Arg Thr Val Asn Glu Asn Phe Leu Ser 65 70 75 80

Leu Gln Leu Asp Pro Ser Ile Ile His Asp Gly Trp Leu Asp Phe Leu 85 90 95

Ser Ser Lys Arg Leu Val Thr Leu Ala Arg Gly Leu Ser Pro Ala Phe 100 105 110

Leu Arg Phe Gly Gly Lys Arg Thr Asp Phe Leu Gln Phe Gln Asn Leu 115 120 125

Arg Asn Pro Ala Lys Ser Arg Gly Gly Pro Gly Pro Asp Tyr Tyr Leu 130 135 140

Lys Asn Tyr Glu Asp Asp Ile Val Arg Ser Asp Val Ala Leu Asp Lys 145 150 155 160

Gln Lys Gly Cys Lys Ile Ala Gln His Pro Asp Val Met Leu Glu Leu 165 170 175

Gln Arg Glu Lys Ala Ala Gln Met His Leu Val Leu Leu Lys Glu Gln 180 185 190

Phe Ser Asn Thr Tyr Ser Asn Leu Ile Leu Thr Ala Arg Ser Leu Asp 195 200 205

Lys Leu Tyr Asn Phe Ala Asp Cys Ser Gly Leu His Leu Ile Phe Ala 210 220

Leu Asn Ala Leu Arg Arg Asn Pro Asn Asn Ser Trp Asn Ser Ser Ser 225 230 235 240

Ala Leu Ser Leu Leu Lys Tyr Ser Ala Ser Lys Lys Tyr Asn Ile Ser 245 250 255

Trp Glu Leu Gly Asn Glu Pro Asn Asn Tyr Arg Thr Met His Gly Arg
260 265 270

Ala Val Asn Gly Ser Gln Leu Gly Lys Asp Tyr Ile Gln Leu Lys Ser 275 280 285 Leu Leu Gln Pro Ile Arg Ile Tyr Ser Arg Ala Ser Leu Tyr Gly Pro 295 Asn Ile Gly Arg Pro Arg Lys Asn Val Ile Ala Leu Leu Asp Gly Phe 315 Met Lys Val Ala Gly Ser Thr Val Asp Ala Val Thr Trp Gln His Cys Tyr Ile Asp Gly Arg Val Val Lys Val Met Asp Phe Leu Lys Thr Arg 345 Leu Leu Asp Thr Leu Ser Asp Gln Ile Arg Lys Ile Gln Lys Val Val Asn Thr Tyr Thr Pro Gly Lys Lys Ile Trp Leu Glu Gly Val Val Thr Thr Ser Ala Gly Gly Thr Asn Asn Leu Ser Asp Ser Tyr Ala Ala Gly 390 Phe Leu Trp Leu Asn Thr Leu Gly Met Leu Ala Asn Gln Gly Ile Asp 410 Val Val Ile Arg His Ser Phe Phe Asp His Gly Tyr Asn His Leu Val 420 425 Asp Gln Asn Phe Asn Pro Leu Pro Asp Tyr Trp Leu Ser Leu Leu Tyr 440 Lys Arg Leu Ile Gly Pro Lys Val Leu Ala Val His Val Ala Gly Leu 455 Gln Arg Lys Pro Arg Pro Gly Arg Val Ile Arg Asp Lys Leu Arg Ile 470 Tyr Ala His Cys Thr Asn His His Asn His Asn Tyr Val Arg Gly Ser 485 490 Ile Thr Leu Phe Ile Ile Asn Leu His Arg Ser Arg Lys Lys Ile Lys 505 Leu Ala Gly Thr Leu Arg Asp Lys Leu Val His Gln Tyr Leu Leu Gln Pro Tyr Gly Gln Glu Gly Leu Lys Ser Lys Ser Val Gln Leu Asn Gly Gln Pro Leu Val Met Val Asp Asp Gly Thr Leu Pro Glu Leu Lys Pro Arg Pro Leu Arg Ala Gly Arg Thr Leu Val Ile Pro Pro Val Thr Met 570

Gly Phe Phe Val Val Lys Asn Val Asn Ala Leu Ala Cys Arg Tyr Arg

585

580

	<211 <212	0 > 1 1 > 1 2 > P 3 > H	12 RT	sapi	ens													
	<400 Asp 1			. Arg	Ser 5	Asp	Val	Ala	Leu	Asp 10	Lys	Gln	Lys	Gly	Cys 15	Lys		
	Ile	Ala	Gln	His 20	Pro	Asp	Val	Met	Leu 25	Glu	Leu	Gln	Arg	Glu 30	Lys	Ala		
	Ala	Gln	Met 35	: His	Leu	Val	Leu	Leu 40	Lys	Glu	Gln	Phe	Ser 45	Asn	Thr	Tyr		
	Ser	Asn 50		ı Ile	Leu	Thr	Ala 55	Arg	Ser	Leu	Asp	Lys 60	Leu	Tyr	Asn	Phe		
	Ala 65	Asp	Cys	s Ser	Gly	Leu 70	His	Leu	Ile	Phe	Ala 75	Leu	Asn	Ala	Leu	Arg 80		
1	Arg	Asn	Pro) Asn	Asn 85	Ser	Trp	Asn	Ser	Ser 90	Ser	Ala	Leu	Ser	Leu 95	Leu		
D 'vi	Lys	Tyr	Ser	Ala		Lys	Lys	Tyr	Asn 105		Ser	Trp	Glu	Leu 110		Asn		
æ	<210	0 > 1	5															
[]	<213	1 > 1 2 > D	.779 NA	sani	ene													
377	<400			sapi	CIID													
				TTTG'	TGCC'	TT C	CCTG	AAGC	CATO	GCC'	rcca	GCA	ACTC	CCG (cccc	CCCGCG	}	60
	TGC	CTAG	CCC	CGGG	GGCT	CT C'	ract:	TGGC	r cto	GTTG	CTCC	ATC	rctc	CCT '	TTCC:	rcccag	1	20
	GCT	GGAG	ACA	GGAG	ACCC'	TT G	CCTG	TAGA	C AG	AGCT	GCAG	GTT	rgaa(GGA A	AAAG	ACCCTG	1	80
	ATTO	CTAC	TTG	ATGT	GAGC	AC C	AAGA	ACCC	A GT	CAGG	ACAG	TCA	ATGA	GAA (CTTC	CTCTCT	2	40
	CTG	CAGC	TGG	ATCC	GTCC	AT C	ATTC	ATGA:	r GG	CTGG	CTCG	ATT	rcct/	AAG (CTCC	AAGCGC	3	00
	TTG	GTGA	CCC	TGGC	CCGG	GG A	CTTT	CGCC	C GC	CTTT	CTGC	GCT	rcgg	GGG (CAAA	AGGACC	3	60
	GAC:	TTCC	TGC	AGTT	CCAG	AA C	CTGA	GGAA	C CC	GGCG	AAAA	GCC	GCGG	GGG (CCCG	GCCCG	4	20
	GAT:	TACT	ATC	TCAA	AAAC'	TA T	GAGG	ATGA	C AT	rgtt	CGAA	GTG	ATGT'	TGC (CTTA	SATAAA	4	80
	CAG	AAAG	GCT	GCAA	GATT	GC C	CAGC	ACCC:	r GA	rgtt	ATGC	TGG	AGCT	CCA I	AAGG	GAGAAG	5	40
	GCA	GCTC	'AGA	TGCA	TCTG	GT T	CTTC'	TAAA	G GA	GCAA'	TTCT	CCA	ATAC'	TTA (CAGT	AATCTC	! 6	00
	ATA	TTAA	CAG	CCAG	GTCT	CT A	GACA	AACT	r TA'	raac'	TTTG	CTG	ATTG	CTC '	TGGA	CTCCAC	! 6	60

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	CTGATA	ATTTG	CTCTAAATGC	ACTGCGTCGT	AATCCCAATA	ACTCCTGGAA	CAGTTCTAGT	720
	GCCCTC	SAGTC	TGTTGAAGTA	CAGCGCCAGC	AAAAAGTACA	ACATTTCTTG	GGAACTGGGT	780
	AATGAC	CCAA	ATAACTATCG	GACCATGCAT	GGCCGGGCAG	TAAATGGCAG	CCAGTTGGGA	840
	AAGGAT	TACA	TCCAGCTGAA	GAGCCTGTTG	CAGCCCATCC	GGATTTATTC	CAGAGCCAGC	900
	TTATA	rggcc	CTAATATTGG	GCGGCCGAGG	AAGAATGTCA	TCGCCCTCCT	AGATGGATTC	960
	ATGAAC	GTGG	CAGGAAGTAC	AGTAGATGCA	GTTACCTGGC	AACATTGCTA	CATTGATGGC	1020
	CGGGTC	GTCA	AGGTGATGGA	CTTCCTGAAA	ACTCGCCTGT	TAGACACACT	CTCTGACCAG	1080
	ATTAGO	BAAAA	TTCAGAAAGT	GGTTAATACA	TACACTCCAG	GAAAGAAGAT	TTGGCTTGAA	1140
	GGTGTC	GTGA	CCACCTCAGC	TGGAGGCACA	AACAATCTAT	CCGATTCCTA	TGCTGCAGGA	1200
	TTCTT	ATGGT	TGAACACTTT	AGGAATGCTG	GCCAATCAGG	GCATTGATGT	CGTGATACGG	1260
	CACTC	TTTT	TTGACCATGG	ATACAATCAC	CTCGTGGACC	AGAATTTTAA	CCCATTACCA	1320
r=1	GACTA	CTGGC	TCTCTCTCCT	CTACAAGCGC	CTGATCGGCC	CCAAAGTCTT	GGCTGTGCAT	1380
D	GTGGCT	rgggc	TCCAGCGGAA	GCCACGGCCT	GGCCGAGTGA	TCCGGGACAA	ACTAAGGATT	1440
14 14				CCACAACCAC				1500
w	ATCATO	CAACT	TGCATCGATC	AAGAAAGAAA	ATCAAGCTGG	CTGGGACTCT	CAGAGACAAG	1560
	CTGGTT	CACC	AGTACCTGCT	GCAGCCCTAT	GGGCAGGAGG	GCCTAAAGTC	CAAGTCAGTG	1620
3	CAACTO	GAATG	GCCAGCCCTT	AGTGATGGTG	GACGACGGGA	CCCTCCCAGA	ATTGAAGCCC	1680
		CCTTC	GGGCCGGCCG	GACATTGGTC	ATCCCTCCAG	TCACCATGGG	CTTTTTTGTG	1740
Ha at Mark mila		SAATG	TCAATGCTTT	GGCCTGCCGC	TACCGATAA			1779
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	<400> GACATT		GAAGTGATGT	TGCCTTAGAT	AAACAGAAAG	GCTGCAAGAT	TGCCCAGCAC	60
	CCTGAT	TGTTA	TGCTGGAGCT	CCAAAGGGAG	AAGGCAGCTC	AGATGCATCT	GGTTCTTCTA	120
	AAGGAG	CAAT	TCTCCAATAC	TTACAGTAAT	CTCATATTAA	CAGCCAGGTC	TCTAGACAAA	180
	CTTTAT	TAACT	TTGCTGATTG	CTCTGGACTC	CACCTGATAT	TTGCTCTAAA	TGCACTGCGT	240
	CGTAAT	CCCA	ATAACTCCTG	GAACAGTTCT	AGTGCCCTGA	GTCTGTTGAA	GTACAGCGCC	300
	AGCAAA	AAAGT	ACAACATTTC	TTGGGAACTG	GGTAAT			336